Scenario comparisons: How much good can we do?

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Updated from the 18th UK Stata Users’ Group Meeting,
13–14 September, 2012
Original version downloadable from the conference website at
http://ideas.repec.org/s/boc/usug12.html
What are scenario comparisons?

- Applied scientists, especially public health scientists, frequently want to know how much good can be caused by a proposed intervention.
- *For instance*, we might want to estimate how much we could decrease the level of a disease, in a dream scenario where the whole world stopped smoking.
- In statistics, scenarios are different versions of a dataset, with the same variables but different values and/or observations.
- We may want to compare different scenarios applied to the same population (with the same parameters).
- Alternatively, we may want to compare the same scenario between different populations (with different parameters), as when standardizing disease rates to a common distribution of gender and age.
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Existing Stata tools for scenario comparisons

- Brady (1998)[1] introduced the Stata Version 5 package `aflogit` for estimating the population attributable fraction (with confidence limits) for cohort and case–control data.
- This is still used, although it sometimes has problems with the 32-character names used in later Stata versions.
- In Stata Version 11, the `margins` command was added, allowing estimation (with confidence limits) of scenario means of a wide range of quantities.
- In Stata Version 12, the `pwcompare` command was added, together with the `pwcompare` option of `margins`, to estimate pairwise differences between scenario means.
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- The punaf suite of packages (Newson, 2013)[4] can be downloaded from SSC, and includes punaf, punafcc, regpar, margprev, marglmean, and now scenttest.

- They use margins to compute confidence intervals for scenario means and proportions and/or their comparisons (differences and ratios), including population attributable (and unattributable) risks and fractions.

- These are estimated (using nlcom) with asymmetric confidence limits, calculated from appropriate Normalizing and variance–stabilizing transformations.

- The results may be saved as estimation results, and/or listed, and/or saved in output datasets, using the SSC package parmest.
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Packages of the punaf suite

These estimate and/or compare marginal means and/or prevalences for one and/or two scenarios (“Scenario 1” and “Scenario 0”), using Normalizing and variance–stabilizing transformations.

<table>
<thead>
<tr>
<th>Package</th>
<th>Computes confidence intervals for:</th>
<th>Transformation</th>
</tr>
</thead>
<tbody>
<tr>
<td>margprev</td>
<td>Marginal prevalences</td>
<td>Logit</td>
</tr>
<tr>
<td>marglmean</td>
<td>Marginal arithmetic means</td>
<td>Log</td>
</tr>
<tr>
<td>regpar</td>
<td>Differences between marginal prevalences (population attributable risks (PARs))</td>
<td>Fisher’s $z$</td>
</tr>
<tr>
<td>punaf</td>
<td>Ratios between marginal arithmetic means (population unattributable fractions (PUFs))</td>
<td>Log</td>
</tr>
<tr>
<td>punafcc</td>
<td>Arithmetic mean risk or hazard ratios (case–control or survival PUFs)</td>
<td>Log</td>
</tr>
<tr>
<td>scenttest</td>
<td>Differences between marginal arithmetic means, or between marginal Poisson rates (PARs)</td>
<td>Identity</td>
</tr>
</tbody>
</table>

A population *attributable* fraction (PAF) is estimated by end–point transformation, subtracting the corresponding PUF from 1, as recommended by Greenland and Drescher (1993)[2].

*Scenario comparisons: How much good can we do?*
Examples in the \texttt{lbw} data

- The \texttt{lbw} dataset was discussed by Hosmer, Lemeshow and Klar (1988)[3], and is distributed on-line by Stata Press.
- It has one observation for each of a sample of 189 pregnancies, and data on the birth weight of the baby, and on a list of predictive variables.
- The most interesting of these variables is probably the mother’s smoking status during pregnancy, coded as the binary variable \texttt{smoke}.
- In our examples, we will try to estimate how much good could be done by eliminating smoking (at least during pregnancy).
- This good is measured using scenario prevalences of low birth weight, which is stored in the binary variable \texttt{low} (birth weight below 2500 grams).
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Logistic regression in the lbw data

After loading the lbw data, we fit a logistic regression model of low with respect to smoke and the confounder race (white, black or other):

```
. logit low i.race i.smoke, or vce(robust);
```

Iteration 0: log pseudolikelihood = -117.336
Iteration 1: log pseudolikelihood = -110.10441
Iteration 2: log pseudolikelihood = -109.98749
Iteration 3: log pseudolikelihood = -109.98736
Iteration 4: log pseudolikelihood = -109.98736

Logistic regression

<table>
<thead>
<tr>
<th></th>
<th>Robust</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Odds Ratio Std. Err.</td>
</tr>
<tr>
<td>----------</td>
<td>-------------------</td>
</tr>
<tr>
<td>race</td>
<td></td>
</tr>
<tr>
<td>black</td>
<td>2.956742 1.420439 2.26 0.024 1.153162 7.581175</td>
</tr>
<tr>
<td>other</td>
<td>3.030001 1.187272 2.83 0.005 1.405753 6.530954</td>
</tr>
<tr>
<td>1.smoke</td>
<td>3.052631 1.10296 3.09 0.002 1.503568 6.197631</td>
</tr>
<tr>
<td>_cons</td>
<td>.1587319 .0515235 -5.67 0.000 .0840173 .2998882</td>
</tr>
</tbody>
</table>

We see that maternal smoking trebles the odds of low birth weight.
But how much good can we do?

- Not many people really understand odds ratios.
- An audience of non–mathematicians might prefer to know what difference it would make, if all pregnant mothers stopped smoking.
- The `regpar` package can answer this question, by comparing prevalences of low birth rate under 2 scenarios.
- “Scenario 0” is the world we live in, where some mothers smoke.
- “Scenario 1” is a fantasy world, where no mothers smoke, but the race distribution stays the same.
- The difference between these prevalences is the population attributable risk (PAR).
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Scenario prevalences and the PAR using `regpar`

We execute `regpar`, as follows:

```
regpar, at(smoke=0);
```

Scenario 0: (as observed) _all
Scenario 1: smoke=0

Symmetric confidence intervals for the logit proportions under Scenario 0 and Scenario 1

and for the z-transformed population attributable risk (PAR)

Total number of observations used: 189

| Coef. Std. Err.  z  P>|z| [95% Conf. Interval] |
|-----------------|-----------------|-----|-----|---------------------|
| Scenario_0 | -.789997 .1519305 -5.20 0.000 -1.087775 -.4922187 |
| Scenario_1 | -1.215955 .2051031 -5.93 0.000 -1.61795 -.8139606 |
| PAR | .0837153 .0266196 3.14 0.002 .0315419 .1358887 |

Asymmetric 95% CIs for the untransformed proportions under Scenario 0 and Scenario 1

and for the untransformed population attributable risk (PAR)

<table>
<thead>
<tr>
<th>Estimate Minimum Maximum</th>
</tr>
</thead>
<tbody>
<tr>
<td>Scenario_0  .3121693 .2520374 .379371</td>
</tr>
<tr>
<td>Scenario_1  .228649  .1654878 .3070471</td>
</tr>
<tr>
<td>PAR         .0835203 .0315315 .1350584</td>
</tr>
</tbody>
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We see that 3.2 to 13.5 percent of all babies might be saved from low birth weight, if no mothers smoked.
Advice for smoking mothers

- Our real aim is to communicate our message to an audience of smoking mothers, and not just to an audience of target-minded public health professionals.
- This non–professional audience might want to know what good they could do for their babies.
- The regpar package can answer this question, too, by comparing prevalences of low birth weight under the 2 scenarios in the subpopulation of smoking mothers.
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**Scenario prevalences and the exposed subpopulation attributable risk for smoking mothers**

We execute `regpar` with the `subpop()` option:

```plaintext
.regpar, at(smoke=0) subpop(if smoke==1);
```

**Scenario comparisons: How much good can we do?**

We see that 7.8 to 34.1 percent of babies *of smoking mothers* might be saved, if none of their mothers smoked.
Returning to our previous audience of target-minded professionals, we might be asked what percent of the “burden” of low birthweight might be removed by eliminating smoking.

The package to answer this is \texttt{punaf}, which calculates population unattributable and attributable fractions.

Prevalences and rates are arithmetic means of non-negative variables.

\texttt{punaf} estimates 2 scenario means of the same non-negative variable, and their ratio, the population unattributable fraction.

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Attributable disease burden as a fraction of the total disease burden

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Prevalences and rates are arithmetic means of non-negative variables.

punaf estimates 2 scenario means of the same non-negative variable, and their ratio, the population unattributable fraction.

punaf then subtracts this ratio from 1 to get the population attributable fraction.
Scenario prevalences and the population unattributable and attributable fractions

We execute `punaf` as follows:

```
punaf, at(smoke=0) eform;
```

Scenario 0: (asobserved) _all
Scenario 1: smoke=0

Confidence intervals for the means under Scenario 0 and Scenario 1 and for the population unattributable faction (PUF)
Total number of observations used: 189

|                | Mean/Ratio | Std. Err. | z   | P>|z| | [95% Conf. Interval] |
|----------------|------------|-----------|-----|------|----------------------|
| Scenario_0     | .3121693   | .0326225  | -11.14 | 0.000 | .2543534 .3831271    |
| Scenario_1     | .228649    | .0361738  | -9.33 | 0.000 | .1676887 .3117704    |
| PUF            | .7324519   | .0818807  | -2.79 | 0.005 | .5883333 .911874     |

95% CI for the population attributable fraction (PAF)

<table>
<thead>
<tr>
<th></th>
<th>Minimum</th>
<th>Maximum</th>
</tr>
</thead>
<tbody>
<tr>
<td>PAF</td>
<td>.2675481</td>
<td>.4116667</td>
</tr>
</tbody>
</table>

The scenario prevalences are estimated, with their ratio, the PUF. This is subtracted from 1 to estimate the PAF, which is 8.8 to 41.2 percent of the Scenario 0 prevalence.
Attributable burden as a fraction of the total burden from smoking mothers

- For completeness, we can estimate the exposed subpopulation attributable fraction.
- This is done using `punaf`, with a `subpop()` option.
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Scenario prevalences and the exposed subpopulation unattributable and attributable fractions

We execute `punaf` as follows:

```
punaf, at(smoke=0) subpop(if smoke==1) eform;
```

| Scenario | Mean/Ratio | Std. Err. | z   | P>|z| | [95% Conf. Interval] |
|----------|------------|-----------|-----|-----|---------------------|
| Scenario_0 | .4054054   | .0572221  | -6.40 | 0.000 | .3074285 .5346073 |
| Scenario_1 | .19209     | .0353824  | -8.96 | 0.000 | .1338801 .2756092 |
| PUF        | .4738221   | .1103706  | -3.21 | 0.001 | .3001505 .7479826 |

95% CI for the population attributable fraction (PAF)

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</thead>
<tbody>
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<td>.5261779</td>
<td>.6998495</td>
</tr>
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</table>

This time, we see that 25.2 to 70.0 percent of the low–weight births from smoking mothers can be attributed to smoking.
Summary: Attributable risks and fractions in the lbw data

- The upper plot gives PARs, measuring prevention in all children.
- The lower plot gives PAFs, measuring prevention in low birth weight children.
- In both cases, proportions prevented are greater for children with smoking mothers.
- These conclusions assume that the association is causal.
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Standardization as out–of–sample prediction

- The punaf suite may also be used to compare the same scenario between different models, as well as vice versa.
- For example, we might fit multiple logit models to multiple independent subpopulation datasets, and then estimate the marginal prevalence that each model would imply in a standard–population dataset, with a standard distribution of gender and age.
- This practice is an example of out–of–sample prediction.
- To do this, packages of the punaf suite all have an option noesample, functioning as the option of the same name for margins.
- We will illustrate this with an example using margprev.
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Skin prick allergy prevalences in the GA\textsuperscript{2}LEN Follow–up Survey

- The GA\textsuperscript{2}LEN Follow-up Survey is part of a multi–centre European study on asthma and allergy.
- Sensitivity to a number of allergens was measured, using skin prick tests, in 13 regional subpopulations distributed over Europe.
- For each allergen in each subpopulation, we fitted a logit model of sensitivity to that allergen in that subpopulation, with respect to age and gender.
- The parameters estimated for each such model were then used to estimate the marginal prevalence expected, if this model applied to a European Standard Population with a standard gender and age distribution.
- This was done using \texttt{margprev}, with the \texttt{noesample} option, in a dataset representing the European Standard Population.
- These standardized marginal prevalences were then compared between regional subpopulations, using a chi–squared heterogeneity test, to detect between–region heterogeneity.
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Logit regression of cat allergy with respect to gender and age in the UK

In the UK subpopulation, we fitted a logit model of cat allergy with respect to gender and age, using sampling–probability weights:

```
. logit spt_cat male fquesagec [pwei=sampwt5], or
```

Iteration 0:  log pseudolikelihood = -1030.8768
Iteration 1:  log pseudolikelihood = -977.80033
Iteration 2:  log pseudolikelihood = -973.41056
Iteration 3:  log pseudolikelihood = -973.39866
Iteration 4:  log pseudolikelihood = -973.39866

Logistic regression

|                       | Odds Ratio | Std. Err. | z    | P>|z|    | [95% Conf. Interval] |
|-----------------------|------------|-----------|------|--------|---------------------|
| spt_cat               |            |           |      |        |                     |
| male                  | 2.527963   | 1.535882  | 1.53 | 0.127  | 0.7684525           |
| fquesagec             | 0.6700974  | 0.2209261 | -1.21| 0.225  | 0.3511585           |
| _cons                 | 0.0794547  | 0.0300632 | -6.69| 0.000  | 0.0378487           |

The parameter \_cons is the odds of cat allergy in 48–year–old women. The others are male–gender and per–decade odds ratios. However, not many people understand those parameters. So...
Dataset representing a European Standard Population

... we now load a dataset with 1 observation per gender per age group, and data on the gender, the minimum and maximum ages in the age group, the mean age (uncentered in years and centered at 48 in decades), and the number in that gender and age group in a European Standard Population:

.use estanpop, clear
.list male agemin agemax agemean fquesagec stanpop, abbr(32) sepby(male)

+--------------------------------------------------------+
<table>
<thead>
<tr>
<th>male  agemin  agemax  agemean  fquesagec  stanpop</th>
</tr>
</thead>
</table>
1. | 0  20  24  22  -2.6  7000 |
2. | 0  25  29  27  -2.1  7000 |
3. | 0  30  34  32  -1.6  7000 |
4. | 0  35  39  37  -1.1  7000 |
5. | 0  40  44  42  -0.6  7000 |
6. | 0  45  49  47  -0.1  7000 |
7. | 0  50  54  52  -0.4  7000 |
8. | 0  55  59  57  -0.9  6000 |

(Here, we have only showed the younger female age groups.)
Cat allergy prevalence standardized to the European Standard Population

Having loaded the dataset, we now use `margprev`, with frequency weights and the `noesample` option, to estimate the marginal odds and prevalence of cat allergy in the European Standard Population, using parameters from the model we fitted earlier for the UK:

```
. margprev [fwei=stanpop], eform noesample
Scenario 1: (asobserved) _all
Confidence interval for the marginal odds
under Scenario 1
Total number of observations used: 134000
```

|                     | Odds  | Std. Err. | z     | P>|z|  | [95% Conf. Interval] |
|---------------------|-------|-----------|-------|------|----------------------|
| Scenario_1          | .1782219 | .07486  | -4.11 | 0.000 | .0782391 .4059742 |

Asymmetric 95% CI for the untransformed marginal prevalence
under Scenario 1

<table>
<thead>
<tr>
<th></th>
<th>Estimate</th>
<th>Minimum</th>
<th>Maximum</th>
</tr>
</thead>
<tbody>
<tr>
<td>Scenario_1</td>
<td>.1512635</td>
<td>.0725619</td>
<td>.2887494</td>
</tr>
</tbody>
</table>

The standardized odds for the UK subpopulation, and for the 12 other regional subpopulations, were input into the SSC package `parmhet` to test for heterogeneity of cat allergy prevalence between European regions. This heterogeneity was detectable ($I^2=46.4\%, \ P=.033$).
References


The original presentation, and the do–file producing the examples in the birth weight data, can be downloaded from the conference website at *http://ideas.repec.org/s/boc/usug12.html*

The packages used in this presentation can be downloaded from SSC, using the `ssc` command.